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FORM PTO-1390	U.S. DEPA	RTMENT OF COMMERCE PATENT AND TRADEMARK OFFICE	ATTORNEY'S DOCKET NUMBER										
TI	RANSMITTAL LETTER	TO THE UNITED STATES	4121-136										
		ED OFFICE (DO/EO/US)	U.S. APPLICATION NO. (If known, see 37 CFR 1.5)										
	CONCERNING A FILIN	G UNDER 35 U.S.C. 371	10/069056										
INTERNATI	ONAL APPLICATION NO.	INTERNATIONAL FILING DATE	PRIORITY DATE CLAIMED										
	PCT/EP00/07 835 07835 11 August 2000 13 August 1999												
TITLE OF INVENTION PARVOVIRUS NS 1 VARIANTS													
APPLICANT	r(s) for do/eo/us	Land											
	, Jurg; ROMMELAERE, . rewith submits to the United States	Designated/Elected Office (DO/EO/US) the follo	owing items and other information:										
2.	This is a FIRST submission of items concerning a filing under 35 U.S.C. 371. This is a SECOND or SUBSEQUENT submission of items concerning a filing under 35 U.S.C. 371. This express request to begin national examination procedures (35 U.S.C. 371(f)) at any time rather than delay examination until the expiration of the applicable time limit set in 35 U.S.C. 371(b) and PCT Articles 22 and 39(1).												
5. 🛭 A a. b. c.	 a. is transmitted herewith (required only if not transmitted by the International Bureau). b. has been transmitted by the International Bureau. 												
6. 🛭 A	translation of the International Ap	plication into English (35 U.S.C. 371(c)(2)).											
7. And a. b. c. d.	are transmitted herewith have been transmitted b have not been made; ho	ernational Application under PCT Article 19 (35) (required only if not transmitted by the Internation transmitted by the International Bureau. Wever, the time limit for making such amendment will not be made.	onal Bureau).										
8. 🗆 A	translation of the amendments to t	he claims under PCT Article 19 (35 U.S.C. 371(c)(3)).										
9. 🛭 A	n oath or declaration of the invento	or(s) (35 U.S.C. 371(c)(4)).*(Unsigned)											
	translation of the annexes to the In 35 U.S.C. 371(c)(5)).	nternational Preliminary Examination Report under	er PCT Article 36										
	16. below concern other document in Information Disclosure Statemer	· ·											
12. 🔲 A	n assignment document for recordi	ing. A separate cover sheet in compliance with 37	7 CFR 3.28 and 3.31 is included.										
. =	FIRST preliminary amendment. SECOND or SUBSEQUENT prel	liminary amendment.											
14. 🔲 A	substitute specification.												
15. 🛭 A	small entity statement.												
16. D O	Other items or information:												

NOTE: This application is being filed with an unsigned Oath or Declaration under the provisions of 37 CFR § 1.53 in order that applicants may secure a filing date of February 11, 2002. Upon receipt of a "Notice to File Missing Parts - Filing Date Granted," an executed Declaration and Power of Attorney, will be filed in the Patent and Trademark Office. The undersigned agent affirmatively states that she has been duly authorized and appointed to file this application on behalf of the applicants and that the Declaration and Power of Attorney to be filed hereafter will confirm the undersigned agent's authorization and appointment. Applicants are entitled to small entity status within the meaning of 37 CFR § 1.9.

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Research Tria	ingle Park, N	IC 27709			
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4121-136 PATENT APPLICATION

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

Jurg Nuesch, et al.

Application No.:

New U.S. National Stage Application of

PCT International Application No. PCT/EP00/07835

International Filing Date:

11 August 2000

Priority Date Claimed:

13 August 1999 (European Appl. No. 99 115 161.4)

U.S. National Phase Filing Date:

Date of mailing identified below

Title:

PARVOVIRUS NS 1 VARIANTS



23448

PATENT TRADEMARK OFFICE

EXPRESS MAIL CERTIFICATE

I hereby certify that I am mailing the attached documents to the Commissioner for Patents on the date specified, in an envelope addressed to the Commissioner for Patents, Box Patent Application, Washington, DC 20231, and Express Mailed under the provisions of 37 CFR 1.10.

Katrina Holland

Name of Person Mailing This Document

Kovina Holan

Signature

February 11, 2002

Date

EV037733273US

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PRELIMINARY AMENDMENT

Commissioner for Patents BOX PATENT APPLICATION Washington, D.C. 20231

Sir:

Prior to examination of the above-identified new national phase patent application, please amend the application, as follows:

In the Specification 1

Please insert on page 1, between the title of the application and the first paragraph, the following new paragraph:

CROSS-REFERENCE TO RELATED APPLICATIONS

This application is filed under the provisions of 35 U. S.C. §371 and claims the priority of International Patent Application No. PCT/EP00/07835 filed August 11, 2000, which in turn claims priority of European Patent Application No. 99 115 161.4 filed August 13, 1999.

On page 4 please amend paragraphs 1, 2 and 3 to read as follows:

According to the invention parvovirus NS1 variants are preferred in which the shift of equilibrium is achieved by mutation of one or several phosphorylation sites. Particularly preferred are parvovirus NS1 variants which have a mutation at one or several of the phosphorylation sites 283, 363, 394 and 463. Even more preferred are the parvovirus NS1 variants S283A (SEQ ID NO. 4), T363A (SEQ ID NO. 6), T394A (SEQ ID NO. 8) and T463A (SEQ ID NO. 10), which are indicated in Table 1 and [figure 1] figures 1.1, 1.2, 1.3 and 1.4. In S283A, a serine is exchanged by an alanine at position 283, in T363A, a threonine is exchanged by alanine at position 363, in T394A a threonine is exchanged by alanine at position 394 and in T463A a threonine is exchanged by alanine at position 463.

Applicants has provided a marked-up version of amended paragraphs and claims 1, 2, 3, 4, 5, 6, 7, 9, 10, 11 and 12 in Appendix A, and a clean set of all pending claims, amended to date, in Appendix B.

A further subject matter of the present invention relates to a nucleic acid, particularly a DNA, which codes for an above parvovirus NS1 variant. Such a DNA comprises preferably:

- (a) the DNA of fig. 1.1 (SEQ ID NO. 3), 1.2 (SEQ ID NO. 5), 1.3 (SEQ ID NO. 7) and 1.4 (SEQ ID NO. 9), respectively
- (b) a DNA hybridizing with the DNA from (a), said DNA comprising the mutated phosphorylation site of the DNA from (a), or
- (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.

The DNA of (a) was deposited with DSMZ (Deutsche Sammlung von ikroorganismen and Zellkulturen) on Aug. 11 1999, i.e. fig. 1.1 as Escherichia coli pRSV-NS: S283A under DSM 12994 (SEQ ID NO. 3), fig. 1.2 as Escherichia coli pRSV-NS: T363A under DSM 12995 (SEQ ID NO. 5), fig. 1.3 as Escherichia coli pRSV-NS: T394A under DSM 12996 (SEQ ID NO: 7) and fig. 1.4 as Escherichia coli pRSV-NS: T463A under DSM 12997 (SEQ ID NO: 9).

On page 7, please amend the last full paragraph to read as follows:

Fig. 1 shows the DNA and amino acid sequences of parvovirus NS1 variants according to the invention (fig. 1.1 (SEQ ID Nos. 3 and 4), 1.2 (SEQ ID Nos. 5 and 6), 1.3 (SEQ ID Nos. 7 and 8) and 1.4 (SEQ ID Nos. 9 and 10)) as compared to parvovirus NS1 wild type (SEQ ID Nos. 1 and 2). In this connection, the mutated sites in the parvovirus Ns 1 variants according to the invention are labeled each.

In the Claims

Please amend claims 1, 2, 3, 4, 5, 6, 7, 9, 10, 11 and 12 to read as follows:

1. A parvovirus NS 1 variant protein having a shifted equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b), wherein the shifted equilibrium is selected from the group consisting of:

- the activities (a) are reduced and eliminated, respectively, and activity (b) is maintained or increased; and
- activity (b) is reduced and eliminated, respectively, and the activities (a) are maintained or increased.
- 2. The parvovirus NS 1 variant protein according to claim 1, wherein one or several phosphorylation sites are mutated.
- 3. The parvovirus NS 1 variant protein according to claim 2, wherein the mutations are located at an amino acid residue site selected from the group consisting of: 283, 363, 394 and/or 463.
- 4. The parvovirus NS 1 variant protein according to claim 3, namely the NS 1 variants S283A (SEQ ID NO. 4), T363A (SEQ ID NO. 6), T394A (SEQ ID NO. 8), and T463A (SEQ ID NO. 10).
- 5. A DNA, coding for the parvovirus NS1 variant protein according to claim 1.
- 6. The DNA according to claim 5, wherein the DNA comprises a member selected from the group consisting of:
 - (a) the DNA of SEQ ID Nos 3, 5, 7 and 9, said DNA comprising a mutated phosphorylation site,
 - (b) a DNA hybridizing with the DNA from (a) under high stringency conditions, said DNA comprising the mutated phosphorylation site of the DNA from (a), or (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.
- 7. An expression vector, comprising the DNA according to claim 6.

- A method of producing the parvovirus NS 1 variant protein according to claim 1, comprising the culturing of the transformant according to claim 8 under suitable conditions.
- 10. An antibody, directed against the parvovirus NS 1 variant protein according to claim 1.
- 11. A Kit comprising at least one member selected from the group consisting of:
 - (a) a parvovirus NS 1 variant protein according to claim 1,
 - (b) a DNA according to claim 5,
 - (c) an antibody according to claim 10, and
 - (d) conventional auxiliary agents, such as solvents, buffers, carriers markers and controls.
- 12. Use of the parvovirus NS 1 variant protein according to claim 1 as a toxin for treating tumoral diseases.

Please add claims 14-18.

- 14. The parvovirus NS 1 variant protein_according to claim 3, wherein the NS 1 variants has a mutation at sites S283A(SEQ ID NO. 4), T363A (SEQ ID NO. 6), T394A (SEQ ID NO. 8), and T463A (SEQ ID NO. 10).
- 15. A DNA, coding for the parvovirus NS1 variant protein according to claim 4.
- 16. An expression vector, comprising the DNA according to claim 5.
- 17. A method of producing the parvovirus NS 1 variant protein according to claim 4, comprising the culturing of the transformant according to claim 8 under suitable conditions.

18. An antibody, directed against the parvovirus NS 1 variant protein according to claim 4.

REMARKS

A marked-up version of amended paragraphs and claims 1, 2, 3, 4, 5, 6, 7, 9, 10, 11 and 12 is included herewith in Appendix A and a clean copy of all pending claims is included in Appendix B.

It is requested that the examination and prosecution of this application proceed on the basis of these amended claims 1-18.

Respectfully submitted,

Marianne Fuierer

Registration No. 39,983

Attorney for Applicants

INTELLECTUAL PROPERTY/ TECHNOLOGY LAW P. O. Box 14329 Research Triangle Park, NC 27709 Phone: (919) 419-9350 Exp. (919) 410-9354

APPENDIX A

In the Specification

Please insert on page 1, between the title of the application and the first paragraph, the following new paragraph:

CROSS-REFERENCE TO RELATED APPLICATIONS.

This application is filed under the provisions of 35 U. S.C. §371 and claims the priority of International Patent Application No. PCT/EP00/ 07835 filed August 11, 2000, which in turn claims priority of European Patent Application No. 99 115 161.4 filed August 13, 1999.

On page 4 please amend paragraphs 1, 2 and 3 to read as follows:

According to the invention parvovirus NS1 variants are preferred in which the shift of equilibrium is achieved by mutation of one or several phosphorylation sites. Particularly preferred are parvovirus NS1 variants which have a mutation at one or several of the phosphorylation sites 283, 363, 394 and 463. Even more preferred are the parvovirus NS1 variants S283A (SEQ ID NO. 4), T363A (SEQ ID NO. 6), T394A (SEQ ID NO. 8) and T463A (SEQ ID NO. 10), which are indicated in Table 1 and [figure 1] figures 1.1, 1.2, 1.3 and 1.4. In_S283A, a serine is exchanged by an alanine at position 283, in T363A, a threonine is exchanged by alanine at position 363, in T394A a threonine is exchanged by alanine at position 394 and in T463A a threonine is exchanged by alanine at position 463.

A further subject matter of the present invention relates to a nucleic acid, particularly a DNA, which codes for an above parvovirus NS1 variant. Such a DNA comprises preferably:

- (a) the DNA of fig. 1.1 (SEQ ID NO. 3), 1.2 (SEQ ID NO. 5), 1.3 (SEQ ID NO. 7) and 1.4 (SEQ ID NO. 9), respectively
- (b) a DNA hybridizing with the DNA from (a), said DNA comprising the mutated phosphorylation site of the DNA from (a), or
- (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.

The DNA of (a) was deposited with DSMZ (Deutsche Sammlung von ikroorganismen and Zellkulturen) on Aug. 11 1999, i.e. fig. 1.1 as Escherichia coli pRSV-NS: S283A under DSM 12994 (SEQ ID NO. 3), fig. 1.2 as Escherichia coli pRSV-NS: T363A under DSM 12995 (SEQ ID NO. 5), fig. 1.3 as Escherichia coli pRSV-NS: T394A under DSM 12996 (SEQ ID NO: 7) and fig. 1.4 as Escherichia coli pRSV-NS: T463A under DSM 12997 (SEQ ID NO. 9).

On page 7, please amend the last full paragraph to read as follows:

Fig. 1 shows the DNA and amino acid sequences of parvovirus NS1 variants according to the invention (fig. 1.1 (SEQ ID Nos. 3 and 4), 1.2 (SEQ ID Nos. 5 and 6), 1.3 (SEQ ID Nos. 7 and 8) and 1.4 (SEQ ID Nos. 9 and 10)) as compared to parvovirus NS1 wild type (SEQ ID Nos. 1 and 2). In this connection, the mutated sites in the parvovirus Ns 1 variants according to the invention are labeled each.

In the Claims

Please amend claims 1, 2, 3, 4, 5, 6, 7, 9, 10, 11 and 12 to read as follows:

- 1. A parvovirus NS 1 variant <u>protein</u> having a shifted equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b), wherein the shifted equilibrium is selected from the group consisting of:
 - the activities (a) are reduced and eliminated, respectively, and activity (b) is maintained or increased; and [or]

- activity (b) is reduced and eliminated, respectively, and the activities (a) are maintained or increased.
- 2. The parvovirus NS 1 variant <u>protein</u> according to claim 1, wherein one or several phosphorylation sites are mutated.
- 3. The parvovirus NS 1 variant <u>protein</u> according to claim 2, wherein the mutations are located at <u>an amino acid residue site</u> [sites] <u>selected from the group consisting</u> of: 283, 363, 394 and/or 463.
- 4. The parvovirus NS 1 variant <u>protein</u> according to claim [2 or]3, namely the NS 1 variants S283A (SEQ ID NO. 4), T363A (SEQ ID NO. 6), T394A (SEQ ID NO. 8), and T463A (SEQ ID NO. 10).
- 5. A DNA, coding for the parvovirus NS1 variant <u>protein</u> according to <u>claim 1</u>. [any one of claims 1 to 4.]
- 6. The DNA according to claim 5, wherein the DNA comprises a member selected from the group consisting of:
 - (a) the DNA of [figure 1] <u>SEQ ID Nos 3, 5, 7 and 9, said DNA comprising a mutated phosphorylation site,</u>
 - (b) a DNA hybridizing with the DNA from (a) <u>under high stringency conditions</u>, said DNA comprising the mutated phosphorylation site of the DNA from (a), or (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.
- 7. An expression vector, comprising the DNA according to claim [5 or] 6.
- 9. A method of producing the parvovirus NS 1 variant <u>protein</u> according to <u>claim 1</u> [any one of claims 1 to 4], comprising the culturing of the transformant according to claim 8 under suitable conditions.

- 10. An antibody, directed against the parvovirus NS 1 variant <u>protein</u> according to <u>claim 1</u> [any one of claims 1 to 4].
- 11. A Kit comprising at least one member selected from the group consisting of:
 - (a) a parvovirus NS 1 variant protein according to claim 1[the invention],
 - (b) a DNA according to claim 5 [the invention, e.g. an expression vector, particularly a parvovirus],
 - (c) an antibody according to claim 10, [the invention, as well as] and
 - (d) conventional auxiliary agents, such as solvents, buffers, carriers markers and controls[,

[wherein of components (a) to (d) one or more representatives can be present each].

12. Use of the parvovirus NS 1 variant <u>protein</u> according to claim 1 [one of claims 1 to 4] as a toxin for treating tumoral diseases.

APPENDIX B

In the Specification

Please insert on page 1, between the title of the application and the first paragraph, the following new paragraph:

CROSS-REFERENCE TO RELATED APPLICATIONS

This application is filed under the provisions of 35 U. S.C. §371 and claims the priority of International Patent Application No. PCT/EP00/07835 filed August 11, 2000, which in turn claims priority of European Patent Application No. 99 115 161.4 filed August 13, 1999.

On page 4 please amend paragraphs 1, 2 and 3 to read as follows:

According to the invention parvovirus NS1 variants are preferred in which the shift of equilibrium is achieved by mutation of one or several phosphorylation sites. Particularly preferred are parvovirus NS1 variants which have a mutation at one or several of the phosphorylation sites 283, 363, 394 and 463. Even more preferred are the parvovirus NS1 variants S283A (SEQ ID NO. 4), T363A (SEQ ID NO. 6), T394A (SEQ ID NO. 8) and T463A (SEQ ID NO. 10), which are indicated in Table 1 and figures 1.1, 1.2, 1.3 and 1.4. In S283A, a serine is exchanged by an alanine at position 283, in T363A, a threonine is exchanged by alanine at position 364 and in T463A a threonine is exchanged by alanine at position 394 and in T463A a threonine is exchanged by alanine at position 463.

A further subject matter of the present invention relates to a nucleic acid, particularly a DNA, which codes for an above parvovirus NS1 variant. Such a DNA comprises preferably:

- (a) the DNA of fig. 1.1 (SEQ ID NO. 3), 1.2 (SEQ ID NO. 5), 1.3 (SEQ ID NO. 7) and 1.4 (SEQ ID NO. 9), respectively
- (b) a DNA hybridizing with the DNA from (a), said DNA comprising the mutated phosphorylation site of the DNA from (a), or
- (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.

The DNA of (a) was deposited with DSMZ (Deutsche Sammlung von ikroorganismen and Zellkulturen) on Aug. 11 1999, i.e. fig. 1.1 as Escherichia coli pRSV-NS: S283A under DSM 12994 (SEQ ID NO. 3), fig. 1.2 as Escherichia coli pRSV-NS: T363A under DSM 12995 (SEQ ID NO. 5), fig. 1.3 as Escherichia coli pRSV-NS: T394A under DSM 12996 (SEQ ID NO: 7) and fig. 1.4 as Escherichia coli pRSV-NS: T463A under DSM 12997 (SEQ ID NO. 9).

On page 7, please amend the last full paragraph to read as follows:

Fig. 1 shows the DNA and amino acid sequences of parvovirus NS1 variants according to the invention (fig. 1.1 (SEQ ID Nos. 3 and 4), 1.2 (SEQ ID Nos. 5 and 6), 1.3 (SEQ ID Nos. 7 and 8) and 1.4 (SEQ ID Nos. 9 and 10)) as compared to parvovirus NS1 wild type (SEQ ID Nos. 1 and 2). In this connection, the mutated sites in the parvovirus Ns 1 variants according to the invention are labeled each.

In the Claims

- 1. A parvovirus NS 1 variant protein having a shifted equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b), wherein the shifted equilibrium is selected from the group consisting of:
 - the activities (a) are reduced and eliminated, respectively, and activity (b) is maintained or increased; and
 - activity (b) is reduced and eliminated, respectively, and the activities (a) are maintained or increased.

- 2. The parvovirus NS 1 variant protein according to claim 1, wherein one or several phosphorylation sites are mutated.
- 3. The parvovirus NS 1 variant protein according to claim 2, wherein the mutations are located at an amino acid residue site selected from the group consisting of: 283, 363, 394 and/or 463.
- 4. The parvovirus NS 1 variant protein according to claim 3, namely the NS 1 variants S283A (SEQ ID NO. 4), T363A (SEQ ID NO. 6), T394A (SEQ ID NO. 8), and T463A (SEQ ID NO. 10).
- 5. A DNA, coding for the parvovirus NS1 variant protein according to claim 1.
- 6. The DNA according to claim 5, wherein the DNA comprises a member selected from the group consisting of :
 - (a) the DNA of SEQ ID Nos 3, 5, 7 and 9, said DNA comprising a mutated phosphorylation site,
 - (b) a DNA hybridizing with the DNA from (a) under high stringency conditions, said DNA comprising the mutated phosphorylation site of the DNA from (a), or (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.
- 7. An expression vector, comprising the DNA according to claim 6.
- 8. A transformant, containing the expression vector according to claim 7.
- 9. A method of producing the parvovirus NS 1 variant protein according to claim 1, comprising the culturing of the transformant according to claim 8 under suitable conditions.

- 10. An antibody, directed against the parvovirus NS 1 variant protein according to claim 4.
- 11. A Kit comprising at least one member selected from the group consisting of:
 - (a) a parvovirus NS 1 variant protein according to claim 4,
 - (b) a DNA according to claim 5,
 - (c) an antibody according to claim 10, and
 - (d) conventional auxiliary agents, such as solvents, buffers, carriers markers and controls.
- 12. Use of the parvovirus NS 1 variant protein according to claim 1 as a toxin for treating tumoral diseases.
- 13. Use of the DNA according to claim 7 as a vector for gene therapy.
- 14. The parvovirus NS 1 variant protein_according to claim 3, wherein the NS 1 variants has a mutation at sites S283A (SEQ ID NO. 4), T363A (SEQ ID NO. 6), T394A (SEQ ID NO. 8), and T463A (SEQ ID NO. 10).
- 15. A DNA, coding for the parvovirus NS1 variant protein according to claim 4.
- 16. An expression vector, comprising the DNA according to claim 5.
- 17. A method of producing the parvovirus NS 1 variant protein according to claim 4, comprising the culturing of the transformant according to claim 8 under suitable conditions.
- 18. An antibody, directed against the parvovirus NS 1 variant protein according to claim 4.

HUBER & SCHUESSLER PATE PATE PATE 10 1 1 FEB 2002

1

Parvovirus NS1 Variants

The present invention relates to parvovirus NS1 variants, DNAs coding for them and methods of producing the parvovirus NS1 variants. Furthermore, this invention concerns antibodies directed against the parvovirus NS1 variants as well as the use of the DNAs and the parvovirus NS1 variants.

Parvovirus designates a genus of the virus family Parvoviridae. The parvovirus genus comprises a number of small, icosaedric viruses that can replicate in the absence of a helper virus. Parvovirus contains a single-stranded DNA having a length of about 5.000 bp. At the 3' and 5' ends of the DNA there is one palindromic sequence each. The DNA codes for two capsid proteins, VP1 and VP2, as well as for two regulatory non-structure proteins, NS-1 and NS-2. The latter proteins are phosphorylated and show nuclear or both cytoplasmic and nuclear localization, respectively. NS1 is necessary for viral DNA replication and participates in the regulation of viral gene expression. Particularly, NS1 transactivates the promoter P38 and exhibits DNA-binding, helicase and DNA-nicking activities. Furthermore, NS1 induces cytotoxic and/or cytostatic stress in sensitive host cells.

Parvoviruses are usually well-tolerated by populations of their natural host, in which they persist without apparent pathological signs. This is due to both the protection of foetuses and neonates by maternal immunity, and the striking restriction of parvovirus replication to a narrow range of target proliferating tissues in adult animals. This host tolerance concerns especially rodent parvoviruses, for example the minute virus of mice (MVM) and H-1 virus in their respective natural hosts, namely mice and rats. In addition, humans can be infected with the latter viruses, without any evidence of

associated deleterious effects from existing epidemiological studies and clinical trials. On the other hand, it is known that certain parvoviruses, and especially rodent parvoviruses, are both oncotropic, i.e. accumulate preferentially in neoplastic versus normal tissues, and oncosuppressive, i.e. have a tumor-suppressive effect towards tumor cells, in various animal models. At least part of the oncosuppressive effect is thought to be due to a direct oncolytic action mediated by NS1. This oncosuppressive effect was also demonstrated against human tumor cells transplanted in recipient animals.

It is considered to use parvoviruses for therapeutic purposes. On the one hand, it seems to be of interest to use parvoviruses as vectors for therapeutic genes, i.e. for introducing such genes into the genome of cells. On the other hand, it is considered to use NS1 of parvoviruses as a toxin for treating tumoral diseases. However, initial experiments showed unsatisfactory results.

Therefore, it is the object of the present invention to provide a product by which parvoviruses and NS1 thereof, respectively, can be used for the above purposes.

According to the invention this is achieved by the subject matters defined in the claims.

The present invention is based on the applicant's findings that it is possible to interfere with the activities of parvovirus NS1 so as to shift the equilibrium existing between the DNA replication and transcription activities (a) and the cytotoxicity activity (b). In particular, he produced parvovirus NS1 variants in which the DNA replication and transcription activities (a) are reduced and eliminated, respectively, whereas the cytotoxicity activity (b) is maintained or raised. Moreover, he produced parvovirus NS1 variants in which the cytotoxicity activity (b) is reduced and eliminated, respectively, whereas the DNA replication and transcription activities (a) are

maintained or raised. Examples of such parvovirus NS1 variants are indicated in Table 1 and figure 1. In addition, the applicant recognized that the above parvovirus NS1 variants and expression vectors coding for them, particularly parvoviruses, respectively, are suitable for therapeutic purposes.

According to the invention, the applicant's findings are used to provide a parvovirus NS1 variant in which the equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b) is shifted.

The expression "parvovirus" comprises any parvovirus, particularly a rodent parvovirus, such as minute virus of mice (MVM) and H-1 virus.

The expression "the equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b) is shifted" refers to the fact that in a parvovirus NS1 variant according to the invention such an equilibrium is shifted as compared to the parvovirus NS1 wild-type. In particular, the equilibrium can be shifted to the effect that the DNA replication and transcription activities (a) are reduced and eliminated, respectively, whereas the cytotoxicity activity (b) is maintained or raised. The cytotoxicity activity (b) can also be reduced and eliminated, respectively, whereas the DNA replication and transcription activities (a) are maintained or raised. Such an equilibrium can be determined by various methods. As regards the determination of the DNA replication activity, reference is made e.g. to methods described in Legendre and Rommelaere, 1992, J. Virol. 66, 5705; Cotmore et al., 1992, Virology 190, 365; Cotmore et al., 1993, J. Virol. 67, 1579, Cotmore and Tattersall, 1994, Embo. J. 13, 4145. As to the determination of the transcription activity reference is made to methods described e.g. in Rhode and Richards, 1987, J. Virol. 61, 2807. Regarding the determination of the cytotoxicity activity reference is made to the below examples.

According to the invention parvovirus NS1 variants are preferred in which the shift of equilibrium is achieved by mutation of one or several phosphorylation sites. Particularly preferred are parvovirus NS1 variants which have a mutation at one or several of the phosphorylation sites 283, 363, 394 and 463. Even more preferred are the parvovirus NS1 variants S283A, T363A, T394A and T463A, which are indicated in Table 1 and figure 1. In S283A, a serine is exchanged by an alanine at position 283, in T394A a threonine is exchanged by alanine at position 363, in T394A a threonine is exchanged by alanine at position 394 and in T463A a threonine is exchanged by alanine at position 394 and in

A further subject matter of the present invention relates to a nucleic acid, particularly a DNA, which codes for an above parvovirus NS1 variant. Such a DNA comprises preferably:

- (a) the DNA of fig. 1.1, 1.2, 1.3 and 1.4, respectively
- (b) a DNA hybridizing with the DNA from (a), said DNA comprising the mutated phosphorylation site of the DNA from (a), or
- (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.

The DNA of (a) was deposited with DSMZ (Deutsche Sammlung von Mikroorganismen and Zellkulturen) on Aug. 11, 1999, i.e. fig. 1.1 as Escherichia coli pRSV-NS:S283A under DSM 12994, fig. 1.2 as Escherichia coli pRSV-NS:T363A under DSM 12995, fig. 1.3. as Escherichia coli pRSV-NS:T394A under DSM 12996 and fig. 1.4 as Escherichia coli pRSV-NS:T463A under DSM 12997.

The expression "hybridizing DNA" refers to a DNA which hybridizes with a DNA from (a) under normal conditions, particularly at 20 (C below the melting point of the DNA. In this connection, the expression "hybridizing" refers to conventional hybridization conditions, preferably to hybridization conditions where 5xSSPE, 1 % SDS, 1xDenhardt's solution are used as solution and the hybridization temperatures are between 35 (C and

70(C, preferably 65(C. The hybridization is followed by a wash step first carried out with 2xSSC, 1 % SDS and then with 0.2xSSC at temperatures between 35(C and 70(C, preferably at 65(C. Furthermore, reference is made to Sambrook et al., Molecular Cloning: A Laboratory Manual, 2nd edition, Cold Spring Harbor Laboratory Press, cold Spring Harbor NY (1989).

A DNA according to the invention can be present in a vector and expression vector, respectively. A person skilled in the art is familiar with examples thereof. In the case of an expression vector for E. coli these are e.g. pGEMEX, pUC derivatives, pGEX-2T, pET3b, T7 based expression vectors and pQE-8. For the expression in yeast, e.g. pY100 and Ycpad1 have to be mentioned while e.g. pKCR, pEFBOS, cDM8, pMSCND, and pCEV4 have to be indicated for the expression in animal cells. The baculovirus expression vector pAcSGHisNT-A is especially suitable for the expression in insect cells.

In a preferred embodiment, the vector containing the DNA according to the invention is a virus, e.g. an adenovirus, vaccinia virus, an AAV virus or a parvovirus, such as MVM or H-1, a parvovirus being preferred. The vector may also be a retrovirus, such as MoMULV, MoMuLV, HaMuSV, MuMTV, RSV or GaLV.

For constructing expression vectors which contain the DNA according to the invention, it is possible to use general methods known in the art. These methods include e.g. in vitro recombination techniques, synthetic methods and in vivo recombination methods as described in Sambrook et al., supra, for example.

Furthermore, the present invention relates to host cells which contain the above described vectors. These host cells include bacteria, yeast, insect and animal cells, preferably mammalian cells. The E. coli strains HB101, DH1, x1776, JM101, JM109, BL21, XL1Blue and SG 13009, the yeast strain Saccharomyces cerevisiae and the animal cells L, A9, 3T3, FM3A, CHO, COS,

Vero, HeLa and the insect cells sf9 are preferred. Methods of transforming these host cells, of phenotypically selecting transformants and of expressing the DNA according to the invention by using the above described vectors are known in the art.

Moreover, the present invention relates to antibodies which specifically recognize an above describe parvovirus NS1 variant, i.e. the region of the parvovirus NS1 variant where the mutation responsible for the shifted equilibrium, particularly a mutated phosphorylation site, is located. The antibodies can be monoclonal, polyclonal or synthetic antibodies or fragments thereof, e.g. Fab, Fv or scFV fragments. Preferably monoclonal antibodies are concerned. For the production it is favorable to immunize animals - particularly rabbits or chickens for a polyclonal antibody and mice for a monoclonal antibody - with an above parvovirus NS1 variant or with fragments thereof. Further boosters of the animals can be effected with the same parvovirus NS1 variant or with fragments thereof. The polyclonal antibody can then be obtained from the animal serum and egg yolk, respectively. The monoclonal antibody can be obtained according to standard methods, reference being made particularly to the method by K÷hler and Milstein (Nature 256 (1975), 495) and Galfrú (Meth. Enzymol. 73 (1981), 3). In this case, mouse myeloma cells are fused with spleen cells originating from the immunized animals. Antibodies according to the invention can be used in many ways, e.g. for the immunoprecipitation of the above described parvovirus NS1 variants or for the isolation thereof. The antibodies can be bound in immunoassays in liquid phase or to a solid carrier. In this connection, the antibodies can be labeled in various ways. The person skilled in the art is familiar with suitable markers and labeling methods. Examples of immunoassays are ELISA and RIA.

The present invention provides parvovirus NS1 variants in which the equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b) is shifted. In particular, parvovirus NS1 variants are provided which have a reduced or no cytotoxicity activity, whereas the DNA replication and transcription activities are maintained or increased. Parvovirus NS1 variants are also provided in which the DNA replication and transcription activities are reduced and eliminated, respectively, whereas the cytotoxicity activity is maintained or raised. Thus, the present invention provides products which are suitable for therapeutic purposes. In particular, expression vectors according to the invention, e.g. parvoviruses, can be used for gene-therapeutic measures. Moreover, parvoviruses NS1 variants according to the invention are suitable as toxins, e.g. for treating tumoral diseases.

Therefore, a kit is also provided for the application of the present invention. This kit comprises the following:

- (a) a parvovirus NS1 variant according to the invention,
- (b) a DNA according to the invention, e.g. an expression vector, particularly a parvovirus,
- (c) an antibody according to the invention, as well as
- (d) conventional auxiliary agents, such as solvents, buffers, carriers, markers and controls.

Of component (a) to (d) one or more representatives can be present each.

Brief description of the drawings

Fig. 1 shows the DNA and amino acid sequences of parvovirus NS1 variants according to the invention (fig. 1.1, 1.2, 1.3 and 1.4) as compared to a parvovirus NS1 wild-type. In this connection, the mutated sites in the parvovirus NS1 variants according to the invention are labeled each.

The present invention is explained by the examples.

Example 1: Preparation and purification of NS1 variants according to the invention

The DNA of the NS1 variant S283A according to the invention was provided as an EcoRV to BstEII fragment obtained by chimeric PCR using two mutagenic primers. This fragment was then inserted into the corresponding cleaved expression vector pTHisNS1 (Nuesch et al., Virology 209, (1995), 122) to obtain pTHis NS1:S283A. Such a vector codes for a fusion protein comprising 6 histidine residues (N terminus partner) and S283A of Fig. 1 (C terminus partner). For expression and purification of S283A the NS1 gene under control of the bacteriophage T7 promoter was transferred into vaccinia virus and expressed in eucaryotic cells by double infection together with vTF7-3 (a vaccinia virus expressing the bacteriophage T7 DNA polymerase). 18 hrs post infection cells were harvested and nuclear extracts prepared. The histidine tagged S283A was then purified by affinity chromatography on Ni-NTA agarose and analyzed by 10 % SDS-PAGE (Nuesch et al., supra).

It showed that a parvovirus NS1 variant according to the invention can be prepared in highly pure form.

The NS1 variants T363A, T394A, and T463A were also produced and purified in the same way.

Example 2: Preparation and detection of an antibody according to the invention

Tubes were coated with purified NS1 variants prepared as in example 1 and monoclonal antibodies (e.g. scFv) specifically binding to S283A were isolated from human synthetic VH+VL scFV phage library (Griffith et al., EMBO J., 13, (1994), 3245) according to standard panning protocols after >5 isolation and amplification procedures. The variable region of the isolated scFv harbored in the phagemid were sequenced to identify NS1

variant interacting partner proteins harboring such binding motifs from comparison with known genes in the gene bank.

It showed that monoclonal antibodies according to the invention can be isolated.

In addition, the NS1 variants were used for immunization of animals in order to obtain poly- or monoclonal antibodies.

Example 3: Characterization of the parvovirus NS1 variants S283A, T363A, T394A and T463A according to the invention

The characterization of the parvovirus NS1 variants comprised the determination of the DNA replication, transcription, cytotoxicity, DNA binding, nicking and helicase activities. Known methods were used for this purpose (cf. description, supra). As regards the determination of the helicase activity reference is made to Stahl et al. 1986, EMBO J. 5, 1999. As to the determination of the nicking activity reference is made to Christensen et al., 1997, J. Virol. 71, 1405 and Nuesch et al., 1995, supra. Regarding the determination of the DNA binding reference is made to Cotmore et al. 1995, J. Virol. 69, 1652. As far as the determination of the cytotoxicity activity is concerned, the following steps were carried out: NS1 variants were transferred into an expression vector containing the NS1 gene under the control of the parvovirus MVMP4 promoter (genuine promoter driving the non-structural genes of MVM), and the green fluorescent protein (EGFP) under control of an additional promoter. These constructs were then transfected into A9 cells using lipofectamine (GibcoBRL) according to the manufacturer's instruction and the impact of the NS1 variant on the viability of the cells tested in time course experiments. Transfected cells were identified by fluorescence of the EGFP. Toxic effects were determined in comparison to wild type NS1 or a vector containing no NS1 gene

as a function of time as well as a measure of cytopathic changes on the cell morphology.

The data indicated in Table 1 were obtained:

Table 1

	S283A	T363A	T394A	T463A	wt
P38-TA	*	-	-	++++	++++
ACCA	+	++++	++	++	++
Nick-1	+	-	-	+++	+++
Nick-2	+++	-	-	++++	++++
Nick-3	++	-	-		++++
Heli	++	-	(+)	++++	++++
Rep	+	-	-	+	++++
Cyto	+++++	++	+++	(+)	+++

Example 4: NS1 variants' expression after transduction using recombinant viral vectors

NS1 expression cassettes containing the NS1 variants according to the invention under control of the parvoviral P4 promoter and a 3'untranslated region from parvovirus MVM to ensure stability and translation of the gene product, were transferred either in a parvovirus genome background as exemplified in example 3, or a heterologous viral genome background, such as vaccinia virus (example 1) or adenovirus. Promoter and terminator regions were exchanged according to the requirements. The nucleic acids containing the NS1 variants were then packaged either in vivo (after transient transfection into eucaryotic cells) or in vitro and the packaged transducing particles were isolated. These transducing units containing NS1 variants were used either for studies concerning gene regulation in tissue culture or animals, but also as therapeutic agents either alone or in combination with other agents (such as cytokines) in gene and cancer therapy

approaches.

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Claims

- A parvovirus NS1 variant having a shifted equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b), wherein
- 5 the activities (a) are reduced and eliminated, respectively, and activity (b) is maintained or increased or
- activity (b) is reduced and eliminated, 10 respectively, and the activities (a) are maintained or increased.
 - 2. The parvovirus NS1 variant according to claim 1, wherein one or several phosphorylation sites are mutated.
 - 3. The parvovirus NS1 variant according to claim 2, wherein the mutations are located at sites 283, 363, 394 and/or 463.
- 20 The parvovirus NS1 variant according to claim 2 or 3, namely the NS1 variants S283A, T363A, T394A, and T463A.
 - 5. A DNA, coding for the parvovirus NS1 variant according to any one of claims 1 to 4.
 - 6. The DNA according to claim 5, wherein the DNA comprises:
 - (a) the DNA of figure 1,
 - (b) a DNA hybridizing with the DNA from (a), said DNA comprising the mutated phosphorylation site of the DNA from (a), or
 - (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.
- 35 7. An expression vector, comprising the DNA according to

claim 5 or 6.

8. A transformant, containing the expression vector according to claim 7.

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A method of producing the parvovirus NS1 variant 9. according to any one of claims 1 to 4, comprising the culturing of the transformant according to claim 8 under suitable conditions.

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- An antibody, directed against the parvovirus NS1 variant 10. according to any one of claims 1 to 4.
- 11. Kit comprising:

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- (a) a parvovirus NS1 variant according to the invention,
- a DNA according to the invention, e.g. an expression vector, particularly a parvovirus,
- (c) an antibody according to the invention, as well as
- conventional auxiliary agents, such as solvents, buffers, carriers, markers and controls,

wherein of components (a) (d) one or more to representatives can be present each.

- 25 Use of the parvovirus NS1 variant according to any one of 12. 1 to 4 as a toxin for treating tumoral diseases.
 - Use of the DNA according to claim 7 as a vector for gene therapy.

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(57) Abstract: The present invention relates to a parvovirus NS1 variant having a shifted equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b). Furthermore, this invention relates to DNAs coding for these parvovirus NS1 variants and methods of producing them. Additionally, this invention concerns antibodies directed against the parvovirus NS1 variants as well as the use of the DNAs and the parvovirus NS1 variants.

/12666 A1

Fig. 1

								W:	ild-	type	NS:	1									2
261	ATGGCTGGAAATGCTTACTCTGATGAAGTTTTGGGAGCAACCAAC														320						
														T							-
321	:			-+-			+				+	-		-+-	-		+			AAAA + TTTT	380
	s		Q											N							-
381				-+-			+				+			-+-		- ~ ~	 +			ACAA + TGTT	440
	D		G		N									D					L		_
441				-+-	- - ~		4				+			-+-		~ ~ ~	+			GGAT + CCTA	50 0
	R	G	A	E	T	Т	W	D	Q	s	E	D	M	E	M	E	T	T	4	D '	-
501		GAAATGACCAAAAAGCAAGTATTCATTTTTGATTCTTTGGTTAAAAAATGTTTATTTGAA CTTTACTGGTTTTTCGTTCATAAGTAAAAACTAAGAAACCAATTTTTTACAAATAAACTT														560					
	E	M	T	K	K	Q	V	F	I	F	מ	s	L	V	ĸ	K	C	L	F	E	-
				-+-	-		+				-			-+ -	- - -		+			ATGG + TACC	620
	٧	L	И	T	K	Ñ	I	F	P	G	D	v	N	M	F	V	Q	H	E	W	-
621				-+-		-	+				+	-	 -	-+-	-	~ = =	+	~		AGCT + ICGA	680
							v							G					Q	A	-
681				-+-	-	-	- - +				+		- ~ -	- + -		- -	+		-	AGCC + ICGG	740
														s							-
	TG1	[AA]	TD.	GC A	ACT	AAC	ACC.	AGC	TGA.	AAG	AAT	TAA	ACT.	AAG:	ΔGA	AAT.	AGC	AGA:	AGA(CAAT	

Fig. 1 (Fortsetzung I)

	AC	ATT	'ACA	CGI	TGA	TTG	TGC	TCC	SACT	TTC	TT	LAT:	TC.	TTC	TCT	TTA	TCG	TCI	TCT	GTTA	
	C	N	v	Q	L	T	P	А	E	R	1	K	\mathbf{r}	R	E	I	A	E	מ	N	~
9.01																				GTGT ÷	960
001																				CACA	
	E	W	V	T	L	L	T	Y	K	H	K	Q	Т	K	K	D	Ÿ	T	K	С	-
861																				TAGT	920
																				ATCA	
	V	L	F	G	N	M	I	A	Y	Y	F	L	T	K	K	K	I	s	T	s	-
921		CCACCAAGAGACGGAGGCTATTTTCTTAGCAGTGACTCTGGCTGG																			
																			GAA		
	P	-	R	-		_	-	_	_		-	_		G					_	L	- ,
981				-+-	-		+	. – – -			+	. 		-+-			+			AACG	1040
	K		G		R	ugu H	AGA L		S		L L		T.	ACT D		GTA M	R R	CGG P	TCT E	TTGC T	
			_	_					_		_	_	_	_	_			-	_	I AAAA	
1041				-+-	-		- - +		~		+		- 	-+-	- 		+	-	- - -	+ 1TTT	1100
		E				T							R		R			T			_
	GA.	AGT	ΤΤC	TAT	TAA	AAC'	TAC	ACT	'TAA	AGA	.GCT	'GGI	'GCA	TAA	AAG	AGT	AAC	CTC	ACC	AGAG	
1101	GAAGTTTCTATTAAAACTACACTTAAAGAGCTGGTGCATAAAAGAGTAACCTCACCAGAG +															1160					
	E	V	S	I	K	T	T	L	ĸ	E	L	V	H	ĸ	R	v	T	Ş	P	E	- -
1161																				AGAA	חמלו
																				CTT	1220
	D	W	M	M	M	Q	P	D	S	Y	I	E	M	М	A	Q	₽	G	G	E	-
1221	AACCTGCTGAAAAATACGCTAGACATTTGTACACTAACTCTAGCCAGAACCAAAACAGCA														1280						
																				rcgt	
																				A	-
1281				-+-			+				+		-	-+-			+				1340
						,														ACTG	
																				D IGCT	
1341				-+-			+				+		~ ~ ~	-÷-	~		+				1400
																				X	_

Fig. 1 (Fortsetzung II)

																				ACCA	3466
1401	TA	AAC	GAC	ACA	AAA	TTT	GTC	TGT	TCC	TCC	gtt	TTC	TTT.	ATG	ACA	AAA	TAA	AGT	ACC	rggt	1460
	I	C	С	V	L	N	R	Q	G	G	K	R	N	T	V	L	F	Н	G	P	~
1461				-+-			+		_ ~ -				-	-+-			+			TGGT + ACCA	1520
							I 							_							~
1521				-+-	~ ~ -		+	- - -	- -	 -	+			-+-			+				1580
	AC	GAT.	ATT.	ACG	TCG	GTT	AÇA	TTT	GAA	AGG	TAA	ATT.	ACT	GAC	ATG	GTT	GTT	CTT	GAA	CTAA	
	С	Y	N	A	A	N	ν	Й	F	P	F	N	D	С	T	N	K	И	L	I	~
1581																				CTCT	1640
ě	AC	CCA'	TCT'	TCT	TCG.	ACC	ATT	gaa	ACC	TGT	CGT	TÇA	TŢŢ	GGT	CAA.	TTA	TCG	GTA	AAC	GAGA	
	W	V	Е	E	A	G	N	F	G	Q	Q	V	N	Q	F	K	A	I	C	S	-
1641																				ACCA	1700
																				rggt	
	G	Q	T	I	R	I	D	Q	ĸ	G	K	G	S	K	Q	I	E	P	T	₽	-
1701		<u> </u>		-+-			+				+~-			-+-		-	+			ACCA + FGGT	1760
							E														~
1761				-+-	~ - - ·		+	- -	- -		+		- .	-+-			4			GCT	1820
	CT.	rgr	GTG/	AGT'	TGG'	TTA	GTC'	TCT	GTC'	ATT	CGA)	AŤT(GTA	AGT/	AGA:	rtg:	rgt/	ATGO	AAE	CGGA	<u>-</u> -
	E	H	Т	Q	P	Ι	R	D	R	M	L	И	I	H	L	T	H	T	L	P	-
1821							TGA												GT2	AAG	1880
	CC																	CAAC	CAI	TTC	
	G	D	F	G	L	V	D	K	N	E	M	P	M	I	C	A	W	L	v	ĸ	~
1881																				TTGG	1940
1001																				ACC	1314
	N	G	¥	Q	s	T	M	A	s	Y	C	A	K	W	G	K	V	P	D	W	~
																				rcgc	2000
1941	AG:	CT	rtt C	SAC	CCG	CCT	CGG:	TTT(CCA	GG:	rtg2	AGG2	ATA:	rrrz rrrz	AAA	ľGA	rcc	AAG	CGI	rgcg	2000
	s	E	N	W	A	E	₽	ĸ	ν	P	T	P	I	N	L	L	G	s	Α	R	-
	TC	ACC2	LTT(CAC	GACZ	ACC	GAA	AAG'	TAC	GCC:	rct(CAG	CAC	JAA(CTA!	rgc	acti	AC:	rccz	ACTT	- A
2001																				+ rcaa	2060

Fig. 1 (Fortsetzung III)

	S	P	P	T	T	P	K,	S	T	P	L	Ş	Q	N	Y	A	L	T	P	L	-
2061	GCATCGGATCTCGAGGACCTGGAGCCTTGGAGCACACCAAATACTCCTGTTGCG															2120					
																				ACGC	
	A	Ş	D	L	Ε	D	Ľ,	A	L	E	P	W	s	T	Þ	N	T	P	V	A	-
2121																				rcaa +	2180
	CCGTGACGTCTTTGGGTCTTGTGACCCCTTCGACCAAGGTTTCGGACGGTTCTACCAGTT																				
	G	T	A	E	T	Q	И	T	G	Ξ	A	G	s	ĸ	A	C	Q	D	G	Q	-
2181																				ACCG	2240
																				rggc	
	L	s	P	T	M	S	Ξ	I	E	Ε	D	L	R	A	C	F	G	A	£	P	-
2241			GAAF CTTI	+			+-				+	-		- 22	279						

Fig. 1.1

1100 - 261 Wildtype-NS1-Sequence

GAAGTTTCTATTAAAACTACACTTAAACAGCTGGTGCATAAAAGAGTAACCTCACCAGAG --+------+-----+----+----+----+----+ 1160 CTTCANAGATAATTTTGATGTGAATTTCTCGACCACGTATTTTCTCATTGGAGTGGTCTC EVSIKTTLKELVHKRVTSPE -→A 5283A

1161 - 2279 Wildtype-NS1-Sequence

Fig. 1.2

1340 - 261 Wildtype-NS1-Sequence

ACAAGAACCTGCAGAATTTTTGCTTTTCATGGCTGGAACTATGTTAAAGTTTGCCATGCT TGTTCITGGACGTCTTAAAAACGAAAAGTACCGACCTTGATACAATTTCAAACGCTACGA TRTCRIFAFHGWNYVKVCHA -→A T363A

1401 - 2279 Wildtype-NS1-Sequence

Fig. 1.3

1400 - 261 Wildtype-NS1-Sequence

ATTTGCTGTGTTTTAAACAGACAAGGAGGCAAAAGAAATACTGTTTTATTTCATGGACCA
1401 ----+----+ 1460
TAAACGACACAAAATTTGTCTGTTCCTCCGTTTTCTTTATGACAAAATAAAGTACCTGGT

ICCVLNRQGGKRNTVLFHGP -

1461 - 2279 Wildtype-NS1-Sequence

Fig. 1.4

1640 - 261 Wildtype-NS1-Sequence

GGTCAAACTATTCGCATTGATCAAAAAGGAAAAGGCAGCAAACAGATTGAACCAACACCA

1641

CCAGTTTGATAAGCGTAACTAGTTTTTCCTTTTCCGTCGTTTGTCTAACTTGGTTGTGGT

G Q T I R I D Q K G K G S K Q I E P T P
A T 463A

1701 - 2279 Wildtype-NS1-Sequence

Residence: In den Wegscheiden1, CH-4132 Muttenz, Switzerland

Post Office Address: Same/

1 Inventor's Signature

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	DECLARATION AND PO FOR PATENT APPLICA	A THE OWNER OF THE PROPERTY OF	ORNEY			ATTORNEY DOCKET NO. 4121-136	
4	As a below named inventor My residence/post office at believe I am the original are listed below) of the sulpar VOVIRUS NS 1 VA the specification of which (X) was filed Fellow Number I hereby state that I have a any amendment(s) referred CFR 1.56. Foreign Application(s) and/or I hereby claim foreign priority by	or, I hereby decaddress and citi, first and sole bject matter when the standard for the standard from	zenship are as starinventor (if only ich is claimed and eto unless the followard and was amendenderstood the corcknowledge the dependent of the corcknowledge the description of the corc description of the corcknowledge the description of the corc description of the corcknowledge the description of the corc	one name is listed for which a pate owing box is checon Serial No. 10/0 ed on	below) on the is sought is sought is sought is sought is sought in the information of 365(b) of	er an original, first and joint inventor (if plural nath on the invention entitled: PCT International Application (if applicable). Ed specification, including the claims, as amended in which is material to patentability as defined from the foreign application(s) for patent or inventor(s) certification.	d by n 37
	365(a) of any PCT international foreign application for patent or	application which inventor(s) certific	designated at least or cate having a filing da	ne country other than te before that of the ap	the United S oplication on	States of America, listed below and have also identified belon which priority is claimed:	v ang
	COUNTRY	APPLICAT	ION NUMBER	DATE FILE	ED	PRIORITY CLAIMED UNDER 35 U.S.C. 119	
	EPO	99 115 161.4		13 August 1999		YES:_X NO:	
9	PCT	PCT/EP00/078	35	11 August 2000		YES: <u>X</u> NO:	
1,	claims of this application is not	disclosed in the property of the discrete discre	ior United States applation as defined in Tit	ication in the manner tle 37, Code of Federa application:	provided by	on(s) listed below and, insofar as the subject matter of each the first paragraph of Title 35, United States Code Section ns, Section 1.56(a) which occurred between the filing date STATUS(patented/pending/abandoned)	112,
	Trademark Office connected the			or agent(s) listed belome. Marianne Fuie		cute this application and transact all business in the Pater o. 39983	it _, and
	Send Correspondence to: Steven J. Hultquist Intellectual Property/Techno P.O. Box 14329 Research Triangle Park, NC					Direct Telephone Calls To: Steven J. Hultquist (919) 419-9350	
^ی اں۔	I hereby declare that all statements that these statements were made of Title 18 of the United States	ents made herein o with the knowled Code and that such	ge that willful false st	atements and the like	so made are	nade on information and belief are believed to be true; and is punishable by fine or imprisonment, or both, under Section the application or any patent issued thereon. Citizenship: Swiss	urthe 100

Date

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	DECLARATION AND POWER OF ATTORNEY : (FOR PATENT APPLICATION (continued)	ATTORNEY DOCKET NO. 4121-136
نر	Full Name of Second Inventor: <u>Jean Rommelaere</u>	Citizenship: (KAY) NAX Belgian
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